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BASE AT POSITIONS 4'X AND 5'X OF DNA SITES USED IN SELECTIONS	AMINO ACIDS SELECTED IN FINGERS 2 AND 3	BINDING SITE SIGNATURE	No. OF CLONES SELECTED AND SCREENED	BASES AT POSITIONS 4'X AND 5'X OF DNA SITES USED IN SELECTIONS	AMINO ACIDS SELECTED IN FINGERS 2 AND 3	BINDING SITE SIGNATURE	No. OF CLONES SELECTED AND SCREENED
4'X 5'X	F2 F3	4'X 5'X	SEQ ID	4'X 5'X	F2 F3	4'X 5'X	SEQ ID
GG	R RLEY *R RSED R RHTH R RSSE R RSSA	GATC GATC	4 20 4 21 1 22 1 23 1 24	CC	T DPHN E HSKS E HRON S DRAN T DRAN	GATC GATC	1 70 4 71 1 72 1 73 2 74
AG	R QVTT *R QSGD R QLAT R QDAH R QRAS K QSTS	ATG GATC	4 25 5 26 2 27 1 28 1 29 1 30	AC	*Y RSQD V QVGH T QLAT	ATG GATC	3 75 2 76 3 77
TG	R SSGD R SASA *R NSGD R LVQN K TGAS R TPSG R TQTA R TSAA	TGATC GATC	1 31 1 32 1 33 1 34 1 35 1 36 1 37 2 38	TA	N TSQA N TSAS K LAQT *E RSRD E RLRD V HLAT V HLTT L VGHM	TGATC GATC	4 78 2 79 1 80 3 81 4 82 1 83 1 84 1 85
CG	R DTSV R DAST R DASA R DTSS	CTGATC GATC	3 39 2 40 1 41 2 42	CA	Y HPAT N HPAN E HMSN L DSRA	CTGATC GATC	2 86 1 87 9 88 2 89
GC	A RNHD *N RSTD *S RSTD S SRHS T RNST T RTST T TRYS T RAQN	GATC GATC	2 43 3 44 6 45 1 46 1 47 1 48 3 49 1 50	GT	*T KSSD *K SSSD *K RSHD A RSSY T RSSS T RKTD	GATC GATC	2 90 1 91 1 92 1 93 1 94 1 95
AC	A QAAT A QGTN *V TSRD S QRGGA S QSTT A TSSS T SSST	ATGATC GATC	4 51 1 52 4 53 1 54 2 55 1 56 1 57	AT	S QIST S QIGA T QYST T QSAS T QSQH T QTSH T QPGH T QDTH K QDST	ATGATC GATC	3 96 1 97 1 98 1 99 1 100 1 101 1 102 1 103 2 104
TC	D TISN N TSTA V TSSL V TSSI V TSNS *R GSND A TTSS T TAGS S TTSS S TSSA S LSTT G LSST	TGATC GATC	1 58 1 59 4 60 1 61 1 62 3 63 1 64 1 65 1 66 1 67 1 68 1 69	TT	T TAST T TASH T TSSV T TSSA S HHTS S HAQT S HATT	TGATC GATC	1 105 1 106 1 107 1 108 1 109 1 110 4 111
				CT	K DHSS T HPST S DSSR	CTGATC GATC	5 112 1 113 1 114

RELATIVE SIGNATURE STRENGTH ■ ≤1.00 ■ 0.80 ■ 0.60 ■ 0.40 ■ 0.20 ■ 0.00

FIG. 3